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; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Synthetic
US-09-638-857-19

Query Match          96.1%; Score 39.4; DB 4; Length 1853;
Best Local Similarity 97.6%; Pred. No. 1.5e-07;
Matches 40; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      1 CCGTTGGTTCCATCACTGCCGTCCCAAAATACAGTTTCCGTC 41
        ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
Db      815 CCGTTGGTTCCATCACTGCCGTCCCAAAATACAGTTTCCGTC 855

RESULT 3
US-08-842-079-16
; Sequence 16, Application US/08842079
; Patent No. 6133434
; GENERAL INFORMATION:
; APPLICANT: BUELL, GARY N.
; APPLICANT: SURPRENANT, ANNMARIE
; APPLICANT: KAWASHIMA, ERIC
; TITLE OF INVENTION: A PURINERGIC RECEPTOR
; FILE REFERENCE: 1430-160
; CURRENT APPLICATION NUMBER: US/08/842,079
; CURRENT FILING DATE: 1997-04-28
; NUMBER OF SEQ ID NOS: 20
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 16
; LENGTH: 3540
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Synthetic
US-08-842-079-16

-Query Match          64.9%; Score 26.6; DB 3; Length 3540;
Best Local Similarity 78.0%; Pred. No. 0.056;
Matches 32; Conservative 0; Mismatches 9; Indels 0; Gaps 0;

QY      1 CCGTTGGTTCCATCACTGCCGTCCCAAAATACAGTTTCCGTC 41
        ||||| ||||| ||||| ||||| ||||| ||||| |||||
Db      910 CAGCTGTGCCATCGTGTCAACCCAAATACAGCTTCCGCC 950

RESULT 4
US-09-638-857-16
; Sequence 16, Application US/09638857
; Patent No. 6509163
; GENERAL INFORMATION:
; APPLICANT: BUELL, GARY N.
; APPLICANT: SURPRENANT, ANNMARIE
; APPLICANT: KAWASHIMA, ERIC
; TITLE OF INVENTION: A PURINERGIC RECEPTOR
; FILE REFERENCE: 1430-160
; CURRENT APPLICATION NUMBER: US/09/638,857
; CURRENT FILING DATE: 2000-08-15
; PRIOR APPLICATION NUMBER: 08/842,079
; PRIOR FILING DATE: 1997-04-28
; NUMBER OF SEQ ID NOS: 20
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 16
; LENGTH: 3540
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Synthetic
US-09-638-857-16

Query Match          64.9%; Score 26.6; DB 4; Length 3540;
Best Local Similarity 78.0%; Pred. No. 0.056;
Matches 32; Conservative 0; Mismatches 9; Indels 0; Gaps 0;
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QY      1 CCGTTGGTTCCATCACTGCCGTCCCAAAATACAGTTTCCGTC 41
        ||||| ||||| ||||| ||||| ||||| ||||| |||||
Db      910 CAGCTGGTCCCATCGTGTCAACCCAAATACAGCTTCCGCC 950

RESULT 5
US-09-328-352-3880
; Sequence 3880, Application US/09328352
; Patent No. 6562958
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
; TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: GTC99-03PA
; CURRENT APPLICATION NUMBER: US/09/328,352
; CURRENT FILING DATE: 1999-06-04
; NUMBER OF SEQ ID NOS: 8252
; SEQ ID NO 3880
; LENGTH: 2223
; TYPE: DNA
; ORGANISM: Acinetobacter baumannii
; US-09-328-352-3880

Query Match          54.6%; Score 22.4; DB 4; Length 2223;
Best Local Similarity 72.5%; Pred. No. 3.1;
Matches 29; Conservative 0; Mismatches 11; Indels 0; Gaps 0;

QY      1 CCGTTGGTTCCATCACTGCCGTCCCAAAATACAGTTTCCGTC 40
        ||||| ||||| ||||| ||||| ||||| ||||| |||||
Db      2171 CAGTTGGCTATATCACTGCCGACCCCAATTGCACCATCTGT 2210

RESULT 6
US-09-181-183-9/c
; Sequence 9, Application US/09181183
; Patent No. 6146866
; GENERAL INFORMATION:
; APPLICANT: VIITANEN, PAUL VEIKKO
; APPLICANT: BACOT, KAREN ONLEY
; APPLICANT: JORDAN, DOUGLAS BRIAN
; TITLE OF INVENTION: LUMAZINE SYNTHASE AND
; TITLE OF INVENTION: RIBOFLAVIN SYNTHASE
; NUMBER OF SEQUENCES: 39
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: E. I. DU PONT DE NEMOURS AND COMPANY
; STREET: 1007 MARKET STREET
; CITY: WILMINGTON
; STATE: DELAWARE
; COUNTRY: UNITED STATES OF AMERICA
; ZIP: 19898
; COMPUTER READABLE FORM:
; MEDIUM TYPE: DISKETTE, 3.50 INCH
; COMPUTER: IBM PC COMPATIBLE
; OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
; SOFTWARE: MICROSOFT WORD VERSION 7.0A
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/181,183
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: FLOYD, LINDA AXAMETHY
; REGISTRATION NUMBER: 33,692
; REFERENCE/DOCKET NUMBER: CL-1083
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 302-992-8112
; TELEFAX: 302-773-0164
; INFORMATION FOR SEQ ID NO: 9:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 627 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
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